Title: METHODS FOR DIAGNOSIS AND TREATMENT OF MDK1 SIGNAL TRANSDUCTION DISORDERS Inventor(s): CIOSSEK ET AL Appl. No.: 08/438,265

FIG. 1A

aagcggccggtctgcagtcggagacttgcaggcagcaaacacggtgcgaacgaa				
10 20 30 M V V Q T R F P S W I I L C Y I W L L G F A H T G E A Q A A ATGGTTGTTCAAACTCGGTTCCCTTCGTGGATTATTTTGTGTTACATCTGGCTGCTTGGCTTTGCACACACGGGGGAGGCGCAGGCTGCG 40 50 60	322			
K E V L L D S K A Q Q T E L E N I S S P P S G M E E I S G AAGGAAGTACTATTACTGGACTCGAAAGCACAAACAGAATTGGAATGGATTTCCTCTCCACCCAGTGGGTGG	412			
70 80 90 L D E N Y TI P I R T Y Q V C Q V M E P N Q N N W L R T N W : TTGGATGAGAACTACACTCCGATAAGAACATACCAGGTGTGCCAGGTCATGGAGCCCAACCAGAACAACTGGCTGCGGACTAACTGGATT	502			
100 S K G N A Q R I F V E L K F T L R D O N S L P G V L G T O K TCTAAAGGCAACGCACAAAGGATTTTTGTAGAATTGAAATTCACCTTGAGGGATTGTAATAGTCTTCCCGGAGTCCTGGGAACTTGCAAG 130 . 140 150	592			
E T F N L Y Y Y E T D Y D T G R N I R E N L Y V K I D T I A GAAACGTTTAATTTGTACTATTATGAAACAGACTACGACACCGCAGGAATATACGAGAAAACCTTTATGTTAAAATAGACACCATTGCT	682			
160 170 180 A D E S F T Q G D L G E R K M K L N T E V R E I G P L S K E GCAGATGAAAGTTTCACACAAGGTGACCCTTGGTGAAAGAAA	772			
190 210. G F Y L A F Q D V G A C I A L V S V K V Y Y K K W T I V E GGATTCTATCTTGCCTTTCAGGATGTAGGGGCTTGCATAGCATTGGTTTCTGTCAAAGTGTACTACAAGAAGTGCTGGACCATTGTTGAG	862			
220 230 240 N L A V F P D T V T G S E F S S L V E V R G T © V S S A E E AACTTAGCTGTCTTTCCAGATACAGTGACTGGTTCGGAATTTTCCTCCTTAGTCGAGGTCCGTGGGACATGTGTCAGCAGTGCCGAGGAA	952			
middlight i dooddighii gait i gairt gallankiaanana aga mahma aga mahma aga aga aga aga aga aga aga aga aga a	1042			
and if divided in the second i	1132			
310 F S D R E G S S R C E C E D G Y Y R A P S D P P Y V A C T R TTCTCTGACCGAGAAGGATCATCCAGGTGGAATGTGAAGATGGGTACTACAGAGCTCCTTCTGATCCACCATACGTTGCATGCA	1222			
340 350 P P S A P Q N L I F N I N Q T T V S L E W S P P A D N G G R CCTCCCTCTGCACCACAGAACCTTATTTTCAATATCAATCA	1312			
370 380 390 N D V T Y R I L © K R © S W E Q G E © V P © G S N I G Y M P AACGATGTCACCTACAGAATACTGTGTAAGCGGTGCAGTTGGGAACAGGGAGAATGTGTGCCATGCGGAAGTAACATTGGATACATGCCC	1402			
420 Q Q T G L E D N Y V T V M D L L A E A N Y T F E V E A V N G CAGCAGACGGGATTAGAGGATAACTATGTCACTGTCATGGACCTACTTGCCCATGCAAATTACACTTTCGAAGTTGAAGCTGTAAATGGA	1492			
V S D L S R S Q R L F A A V S I T T G Q A A P S Q V S G V M GTTTCGGACTTAAGCAGATCCCAGAGGGCTCTTCGCTGCTGTTAGCATCACCACCGGTCAAGCAGCTCCCTCGCAAGTGAGTG	1582			
460 470 480 K E R V L Q R S V Q L S W Q E P E E P N G V I T E Y E I K Y AAGGAGCGAGTACTGCAGCGGAGTATCAAGTATCAAGTAT	1672			
490 510 Y E K D Q R E R T Y S T L K T K S T S A S I N N L K P G T V TATGAGAAAGATCAACGGGAAAGGACGTACTCAACACTCAAAACCAAGTCCACCTCCGCCTCCATTAATAATCTGAAAACCGGGAACAGTG	1762			
520 540 Y V F Q I R A V T A A G T G N Y S P R L D V A T L E E A S G TACGTCTTTCAGATCCGGGCGGTCACTGCTGCCGGTTATGGAAACTACAGCCCTAGGCTTGATGTTGCCACACTTGAGGAAAGCTTCAGGT	1852			

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FIG. 1B 550 560 570 K M F E A T A V S S E Q N P V I I A V V A V A G T I L V AAAATGTTTGAAGCGACAGCAGCATCATCTTGGTG 1	1942
580 FMVFGFIIGRRHCGYSKADOEGDEELYFHF TTCATGGTGTTCGGCTTCATCATTGGAAGAAGGCACTGTGGTTATAGCAAGGCTGACCAAGAAGGGGATGAAGAACTCTACTTTCATITI 2 610 620 630	2032
610 620 630 K F P G T K T Y I D P E T Y E D P N R A V H Q F A K E L D A AAATTTCCAGGCACCAAAACCTACATTGACCCTGAAACCTATGAGGACCCAAATAGAGCTGTCCATCAATTCGCCAAGGAGCTAGATGCC 2 640 650 660	2122
S C I K I E R V I G A G E F G E V C S G R L K L P G Q R D V TCCTGTATTAAAATTGAGCGTGTGATTGGTGCAGGAGAATTTGGAGAAGTTTGCAGTGGTCGTTTGAAACTTCCGGGCCAGAGAGATGTT 2 670 680 690	2212
A V A I K T L K V G Y T E K Q R R D F L C E A S I M G Q F D GCAGTGGCCATAAAAACCCTGAAAGTTGGTTACACAGAAAAGCAAAGGAGGGGCACTTTTTATGCGAAGCAAGC	2302
H P N V V H L E G V V T R G K P V M I V I E F M E N G A L D CACCCAAATGTCGTCCATTTGGAAGGGGTTGTTACAAGAGGGAAGCCTGTCATGATTGTGATAGAGTTCATGGAGAAATGGAGCCCTGGAT 730 740 750	2392
A F L R K H D G Q F T V I Q L V G M L R G I A A G M R Y L A GCATTTCTCAGGAAACACGATGGGCAGTTTACAGTCATTCAGTTGGTAGGAATGTTGAGAGGTATTGCCGCTGCCATGCGATACTTGGCT 760 770 780	2482
D M G Y V H R D L Å A R N I L V N S N L V C K V S D F G L S GATATGGGATACGTTCACAGGGACCTTGCAGCGCGCAACATCCTTGTCAACAGCAATCTTGTTTGT	2572
790 800 810 R V I E D D P E A V Y T T T G G K I P V R W T A P E A I O Y CGGGTTATAGAGGATGATCCCGAAGCTGTCTACACCACGACTGGTGGAAAAATTCCAGTAAGGTGGACTGCACCGGAAGCCATTCAATAC	2662
820 830 840 R K F T S A S D V W S Y G I V M W E V M S Y G E R P Y W D M CGGAAGTTCACCTCAGCCAGCGATGTGTGGAGCTATGGGATTGTCATGTGGGAAGTGATGTCTTATGGAGAAAGACCTTACTGGGACATG 850 870	2752
S N Q D V I K A I E E G Y R L P A P M D C P A G L H Q L M L TCAAATCAAGATGTCATTAAAGCGATAGAAGAAGGTTATCGTTTGCCGGCGCCCATGGATTGCCCAGCTGGTCTTCACCAGCTAATGCTG	2842
880 900 D C W Q K D R A E R P K P E Q I V G I L D K M I R N R S S L GATTGTTGGCAGAAAGATCGGGCGGAAAGGCCAAAGTTTGAGCAGATAGTCGGAATTCTAGACAAAATGATTCGAAACCCAAGTAGTCTG	2932
910 930 K T P L G T C S R P L S P L D O S T P D F T A F C S V G E AAAACACCCCTGGGAACTTGTAGTAGACCCTTAAGCCCTCTTCTGGACCAGAGCACTCCTGACTTCACTGCCTTCTGTTCAGTTGGAGAA	3022
940 950 950 960 W L Q A I K M E R Y K D N F T A A G Y N S L E S V A R M T I TGGTTGCAAGCTATTAAAATGGAAAGGTATAAAGGACAACTTCACAGCAGCAGCGGTTACAACTCACTC	3112
970 980 990 DDVMSLGITLVGEQKKIMSSIQTMRAQMLH GATGATGTGATGAGTTTAGGGATCACACTGGTTGGCCATCAAAAGAAGATCATGAGCAGCATCCAGACTATGCGGGCACAAATGTTGCAT	3202
L H G T G I Q V * TTACACGGAACAGGCATCCAAGTGTGA	3229
atatactgggtccttcctacaagaaagaagattttagaagcacctccagacttgaactcctaagtgccaccagaatatacaaaaaggg aatttaggatccaccactggtggccaggaacacagcagagacaataaacaaagtactacctgaaaaacatcccaacacttgagctctcg	3319 3409 3499 3589 3628
ttgtaaaaatcagataacatactaacttttcaatgaaaaaaaa	3769 3859 3949 4039

FIG. 24

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2002	2092	2182 2272 2362 2452 2542 2632 2722 2812 2901
590 V A V A G T I I L V F M V F G F I I G R R H C G Y S K A D Q GTGGCTGTAGCAGGACCATCATCTTGGTGTTCATTGGAAGAGGCACTGTGGTTATAGCAAGGCTGACCAA 610	E G D ·E E L Y F H S L V T N E H L S V L * GAAGGGGATGAAGAACTCTACTTTCATT <u>CTTTAGTAACAAATGAGCACCTGTCAGTTTTATAA</u> accgcaacaataactgtttaagacaat	caattttggataaacaatcaactacagcagaataaafcaagatttttaagtcccattttcctttatacattctgcttattttgttgttgt atgtttattttttaaactctgatcttgattgaatgtgataccataagcacagttaggctgcagtgtaaatatataaagacattgttctga gagcagtacgattccatggaaagattgtttggtggttttgttaaaattaataaaggatttttaaggatatagtgtaatttcttcattgc attaatataaccaaatatgcctacctatctttgtcttgaaccaaatgaatagatttggaatactttattgtaattgaatttcattga ttgactgagcatttatgtgttacctgcatgcttctgggtgcattgaaatattttaacttttaaaatgatactatgttgtatttgatataaa ctacctttgtgaggcatactggctacctcctctattagctaagatcttccaaagccttataatgaaaaggttatataaaccatttctc tttcaaatcactgtcatactggtcacggatccaggaatattgtaaaattttctaatttactctgcactttgtatatcagcctctatta ccctcaaggtgaatataaaaactatgtcttttgaatatttctctttgattttgtgatagcagtcctcatatttgtaatattatagaaaaaaaa

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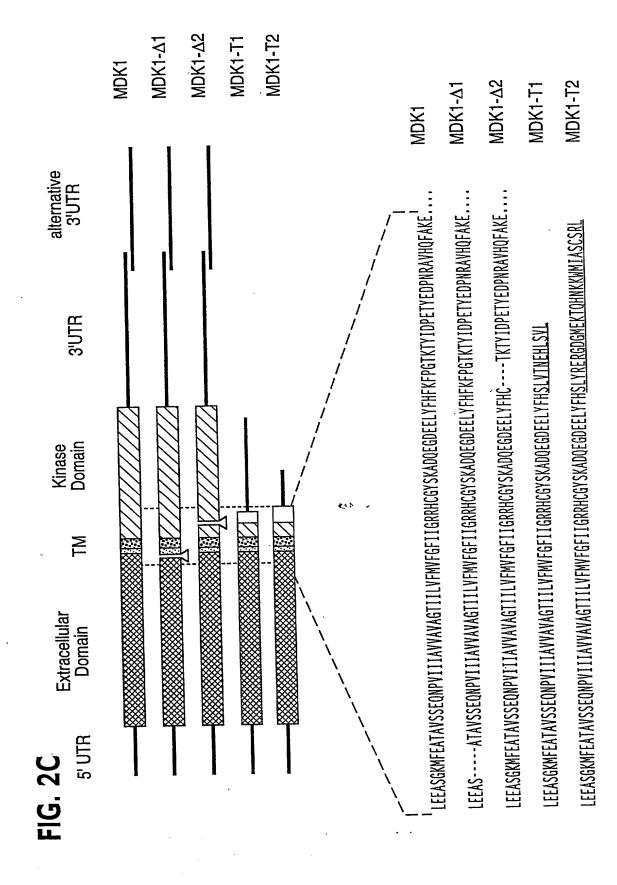
Appl. No.: 08/438,265

FIG. 2B

MDK1-T2

attagcattgtgaacctgactgtaatcctcttccggaaagagatgagatgctattgcgatgagaatgtacaacttgcaccttgaaatc 2272 2323 tttttgataattagtgctcaggggaggggggggggagtagaggaagcaaa

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